

F Harold

1646

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RAW SEQUENCE LISTING  
PATENT APPLICATION US/08/968,800

DATE: 04/15/1999  
TIME: 11:14:59

Input Set: H968800.RAW

This Raw Listing contains the General Information  
Section and up to first 5 pages.

1 <110> APPLICANT: Hyseq, Inc.  
2 <120> TITLE OF INVENTION: A NOVEL EGF MOTIF PROTEIN OBTAINED FROM  
3 A cDNA LIBRARY OF FETAL LIVER-SPLEEN  
4 <130> FILE REFERENCE: 20411-720  
5 <140> CURRENT APPLICATION NUMBER: US/08/968,800  
6 <141> CURRENT FILING DATE: 1997-11-22  
7 <160> NUMBER OF SEQ ID NOS: 20  
8 <170> SOFTWARE: FastSEQ for Windows Version 3.0  
9 <210> SEQ ID NO 1  
10 <211> LENGTH: 300  
11 <212> TYPE: DNA  
12 <213> ORGANISM: Homo sapiens  
13 <220> FEATURE:  
14 <221> NAME/KEY: CDS  
15 <222> LOCATION: (1)...(300)  
16 <221> NAME/KEY: misc\_feature  
17 <222> LOCATION: (1)...(300)  
18 <223> OTHER INFORMATION: n = A,T,C or G  
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22 1 5 10 15  
23 gga tgt aag ttt ggt gag tgc gtg gga cca aac aaa tgc aga tgc ttt 96  
24 Gly Cys Lys Phe Gly Glu Cys Val Gly Pro Asn Lys Cys Arg Cys Phe  
25 20 25 30  
26 cca gga tac acc ggg aaa acc tgc agt caa gat gtg aat gag tgt gga 144  
27 Pro Gly Tyr Thr Gly Lys Thr Cys Ser Gln Asp Val Asn Glu Cys Gly  
28 35 40 45  
29 atg aaa ccc cgg cca tgc caa cac aga tgt gtg aat aca cac gga agc 192  
30 Met Lys Pro Arg Pro Cys Gln His Arg Cys Val Asn Thr His Gly Ser  
31 50 55 60  
32 tac aag tgc ttt tgc ctc agt ggc cac atg ctc atg cca gat gct acg 240  
33 Tyr Lys Cys Phe Cys Leu Ser Gly His Met Leu Met Pro Asp Ala Thr  
34 65 70 75 80  
35 tgt gtg aac tcn agg aca tgt gcc atg ata aac tgt cag tat agc tgt 288  
36 Cys Val Asn Xaa Arg Thr Cys Ala Met Ile Asn Cys Gln Tyr Ser Cys  
37 85 90 95  
38 gaa gac aca gaa 300  
39 Glu Asp Thr Glu  
40 100  
41 <210> SEQ ID NO 2  
42 <211> LENGTH: 1611  
43 <212> TYPE: DNA  
44 <213> ORGANISM: Homo sapiens

P/S  
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RAW SEQUENCE LISTING  
PATENT APPLICATION US/08/968,800DATE: 04/15/1999  
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 46 <221> NAME/KEY: CDS  
 47 <222> LOCATION: (1)...(1506)  
 48 <221> NAME/KEY: misc\_feature  
 49 <222> LOCATION: (1)...(1611)  
 50 <223> OTHER INFORMATION: n = A,T,C or G  
 51 <400> SEQUENCE: 2

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1 5 10 15	
gga tgt aag ttt ggt gag tgc gtg gga cca aac aaa tgc aga tgc ttt	96
Gly Cys Lys Phe Gly Glu Cys Val Gly Pro Asn Lys Cys Arg Cys Phe	
20 25 30	
cca gga tac acc ggg aaa acc tgc agt caa gat gtg aat gag tgt gga	144
Pro Gly Tyr Thr Gly Lys Thr Cys Ser Gln Asp Val Asn Glu Cys Gly	
35 40 45	
atg aaa ccc cgg cca tgc caa cac aga tgt gtg aat aca cac gga agc	192
Met Lys Pro Arg Pro Cys Gln His Arg Cys Val Asn Thr His Gly Ser	
50 55 60	
tac aag tgc ttt tgc ctc agt ggc cac atg ctc atg cca gat gct acg	240
Tyr Lys Cys Phe Cys Leu Ser Gly His Met Leu Met Pro Asp Ala Thr	
65 70 75 80	
tgt gtg aac tcn agg aca tgt gcc atg ata aac tgt cag tat agc tgt	288
Cys Val Asn Xaa Arg Thr Cys Ala Met Ile Asn Cys Gln Tyr Ser Cys	
85 90 95	
gaa gac aca gaa gaa ggg cca cag tgc ctg tgt cca tcc tca gga ctc	336
Glu Asp Thr Glu Glu Gly Pro Gln Cys Leu Cys Pro Ser Ser Gly Leu	
100 105 110	
cgc ctg gcc cca aat gga aga gac tgt cta gat att gat gaa tgt gcc	384
Arg Leu Ala Pro Asn Gly Arg Asp Cys Leu Asp Ile Asp Glu Cys Ala	
115 120 125	
tct ggt aaa gtc atc tgt ccc tac aat cga aga tgt gtg aac aca ttt	432
Ser Gly Lys Val Ile Cys Pro Tyr Asn Arg Arg Cys Val Asn Thr Phe	
130 135 140	
gga agc tac tac tgc aaa tgt cac att ggt ttc gaa ctg caa tat atc	480
Gly Ser Tyr Tyr Cys Lys Cys His Ile Gly Phe Glu Leu Gln Tyr Ile	
145 150 155 160	
agt gga cga tat gac tgt ata gat ata aat gaa tgt act atg gat agc	528
Ser Gly Arg Tyr Asp Cys Ile Asp Ile Asn Glu Cys Thr Met Asp Ser	
165 170 175	
cat acg tgc agc cac cat gcc aat tgc ttc aat acc caa ggg tcc ttc	576
His Thr Cys Ser His His Ala Asn Cys Phe Asn Thr Gln Gly Ser Phe	
180 185 190	
aag tgt aaa tgc aag cag gga tat aaa ggc aat gga ctt cgg tgt tct	624
Lys Cys Lys Cys Lys Gln Gly Tyr Lys Gly Asn Gly Leu Arg Cys Ser	
195 200 205	
gct atc cct gaa aat tct gtg aag gaa gtc ctc aga gca cct ggt acc	672
Ala Ile Pro Glu Asn Ser Val Lys Glu Val Leu Arg Ala Pro Gly Thr	
210 215 220	
atc aaa gac aga atc aag aag ttg ctt gct cac aaa aac agc atg aaa	720

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**RAW SEQUENCE LISTING**  
**PATENT APPLICATION US/08/968,800**
**DATE: 04/15/1999**  
**TIME: 11:14:59**
**Input Set: H968800.RAW**

95	Ile Lys Asp Arg Ile Lys Lys Leu Leu Ala His Lys Asn Ser Met Lys			
96	225	230	235	240
97	aag aag gca aaa att aaa aat gtt acc cca gaa ccc acc agg act cct			768
98	Lys Lys Ala Lys Ile Lys Asn Val Thr Pro Glu Pro Thr Arg Thr Pro			
99	245	250	255	
100	acc cct aag gtg aac ttg cag ccc ttc aac tat gaa gag ata gtt tcc			816
101	Thr Pro Lys Val Asn Leu Gln Pro Phe Asn Tyr Glu Glu Ile Val Ser			
102	260	265	270	
103	aga ggc ggg aac tct cat gga ggt aaa aaa ggg aat gaa gag aaa atg			864
104	Arg Gly Gly Asn Ser His Gly Gly Lys Lys Gly Asn Glu Glu Lys Met			
105	275	280	285	
106	aaa gag ggg ctt gag gat gag aaa aga gaa gag aaa gcc ctg aag aat			912
107	Lys Glu Gly Leu Glu Asp Glu Lys Arg Glu Glu Lys Ala Leu Lys Asn			
108	290	295	300	
109	gac ata gag gag cga agc ctg cga gga gat gtg ttt ttc cct aag gtg			960
110	Asp Ile Glu Glu Arg Ser Leu Arg Gly Asp Val Phe Phe Pro Lys Val			
111	305	310	315	320
112	aat gaa gca ggt gaa ttc ggc ctg att ctg gtc caa agg aaa gcg cta			1008
113	Asn Glu Ala Gly Glu Phe Gly Leu Ile Leu Val Gln Arg Lys Ala Leu			
114	325	330	335	
115	act tcc aaa ctg gaa cat aaa gat tta aat atc tcg gtt gac tgc agc			1056
116	Thr Ser Lys Leu Glu His Lys Asp Leu Asn Ile Ser Val Asp Cys Ser			
117	340	345	350	
118	ttc aat cat ggg atc tgt gac tgg aaa cag gat aga gaa gat gat ttt			1104
119	Phe Asn His Gly Ile Cys Asp Trp Lys Gln Asp Arg Glu Asp Asp Phe			
120	355	360	365	
121	gac tgg aat cct gct gat cga gat aat gct att ggc ttc tat atg gca			1152
122	Asp Trp Asn Pro Ala Asp Arg Asp Asn Ala Ile Gly Phe Tyr Met Ala			
123	370	375	380	
124	gtt ccg gcc ttg gca ggt cac atg aaa gac att ggc cga ttg aaa ctt			1200
125	Val Pro Ala Leu Ala Gly His Met Lys Asp Ile Gly Arg Leu Lys Leu			
126	385	390	395	400
127	ctc cta cct gac ctg caa ccc caa agc aac ttc tgt ttg ctc ttt gat			1248
128	Leu Leu Pro Asp Leu Gln Pro Gln Ser Asn Phe Cys Leu Leu Phe Asp			
129	405	410	415	
130	tac cgg ctg gcc gga gac aaa gtc ggg aaa ctt cga gtg ttt gtg aaa			1296
131	Tyr Arg Leu Ala Gly Asp Lys Val Gly Lys Leu Arg Val Phe Val Lys			
132	420	425	430	
133	aac agt aac aat gcc ctg gca tgg gag aag acc acg agt gag gat gaa			1344
134	Asn Ser Asn Asn Ala Leu Ala Trp Glu Lys Thr Thr Ser Glu Asp Glu			
135	435	440	445	
136	aag tgg aag aca ggg aaa att cag ttg tat caa gga act gat gct acc			1392
137	Lys Trp Lys Thr Gly Lys Ile Gln Leu Tyr Gln Gly Thr Asp Ala Thr			
138	450	455	460	
139	aaa agc atc att ttt gaa gca gaa cgt ggc aag ggc aaa acc ggc gaa			1440
140	Lys Ser Ile Ile Phe Glu Ala Glu Arg Gly Lys Gly Lys Thr Gly Glu			
141	465	470	475	480
142	atc gca gtg gat ggc gtc ttg ctt gtt tca ggc tta tgt cca gat agc			1488
143	Ile Ala Val Asp Gly Val Leu Leu Val Ser Gly Leu Cys Pro Asp Ser			
144	485	490	495	

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RAW SEQUENCE LISTING  
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145      ctt tta tct gtg gan nnc tgaatggtagtac tatctttata tttgactttg          1536
146      Leu Leu Ser Val Xaa Xaa
147          500
148      tatgtcagtt ccctgggtttt tttgatattt catcatagga cctctggcat tttaaaattha  1596
149      ctatgtaaaa aatttg          1611
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151 <211> LENGTH: 100
152 <212> TYPE: PRT
153 <213> ORGANISM: Homo sapiens
154 <400> SEQUENCE: 3
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157     Gly Cys Lys Phe Gly Glu Cys Val Gly Pro Asn Lys Cys Arg Cys Phe
158         20          25           30
159     Pro Gly Tyr Thr Gly Lys Thr Cys Ser Gln Asp Val Asn Glu Cys Gly
160         35          40           45
161     Met Lys Pro Arg Pro Cys Gln His Arg Cys Val Asn Thr His Gly Ser
162         50          55           60
163     Tyr Lys Cys Phe Cys Leu Ser Gly His Met Leu Met Pro Asp Ala Thr
164         65          70           75           80
165     Cys Val Asn Ser Arg Thr Cys Ala Met Ile Asn Cys Gln Tyr Ser Cys
166         85          90           95
167     Glu Asp Thr Glu
168         100
169 <210> SEQ ID NO 4
170 <211> LENGTH: 537
171 <212> TYPE: PRT
172 <213> ORGANISM: Homo sapiens
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175 <222> LOCATION: (1)...(537)
176 <223> OTHER INFORMATION: Xaa = Any Amino Acid
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181         20          25           30
182     Pro Gly Tyr Thr Gly Lys Thr Cys Ser Gln Asp Val Asn Glu Cys Gly
183         35          40           45
184     Met Lys Pro Arg Pro Cys Gln His Arg Cys Val Asn Thr His Gly Ser
185         50          55           60
186     Tyr Lys Cys Phe Cys Leu Ser Gly His Met Leu Met Pro Asp Ala Thr
187         65          70           75           80
188     Cys Val Asn Ser Arg Thr Cys Ala Met Ile Asn Cys Gln Tyr Ser Cys
189         85          90           95
190     Glu Asp Thr Glu Glu Gly Pro Gln Cys Leu Cys Pro Ser Ser Gly Leu
191         100         105          110
192     Arg Leu Ala Pro Asn Gly Arg Asp Cys Leu Asp Ile Asp Glu Cys Ala
193         115         120          125
194     Ser Gly Lys Val Ile Cys Pro Tyr Asn Arg Arg Cys Val Asn Thr Phe

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RAW SEQUENCE LISTING  
PATENT APPLICATION US/08/968,800DATE: 04/15/1999  
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195	130	135	140
196	Gly Ser Tyr Tyr Cys Lys Cys His Ile Gly Phe Glu Leu Gln Tyr Ile		
197	145	150	155
198	Ser Gly Arg Tyr Asp Cys Ile Asp Ile Asn Glu Cys Thr Met Asp Ser		160
199	165	170	175
200	His Thr Cys Ser His His Ala Asn Cys Phe Asn Thr Gln Gly Ser Phe		
201	180	185	190
202	Lys Cys Lys Cys Lys Gln Gly Tyr Lys Gly Asn Gly Leu Arg Cys Ser		
203	195	200	205
204	Ala Ile Pro Glu Asn Ser Val Lys Glu Val Leu Arg Ala Pro Gly Thr		
205	210	215	220
206	Ile Lys Asp Arg Ile Lys Lys Leu Leu Ala His Lys Asn Ser Met Lys		
207	225	230	235
208	Lys Lys Ala Lys Ile Lys Asn Val Thr Pro Glu Pro Thr Arg Thr Pro		240
209	245	250	255
210	Thr Pro Lys Val Asn Leu Gln Pro Phe Asn Tyr Glu Glu Ile Val Ser		
211	260	265	270
212	Arg Gly Gly Asn Ser His Gly Gly Lys Gly Asn Glu Glu Lys Met		
213	275	280	285
214	Lys Glu Gly Leu Glu Asp Glu Lys Arg Glu Glu Lys Ala Leu Lys Asn		
215	290	295	300
216	Asp Ile Glu Glu Arg Ser Leu Arg Gly Asp Val Phe Pro Lys Val		
217	305	310	315
218	Asn Glu Ala Gly Glu Phe Gly Leu Ile Leu Val Gln Arg Lys Ala Leu		320
219	325	330	335
220	Thr Ser Lys Leu Glu His Lys Asp Leu Asn Ile Ser Val Asp Cys Ser		
221	340	345	350
222	Phe Asn His Gly Ile Cys Asp Trp Lys Gln Asp Arg Glu Asp Asp Phe		
223	355	360	365
224	Asp Trp Asn Pro Ala Asp Arg Asp Asn Ala Ile Gly Phe Tyr Met Ala		
225	370	375	380
226	Val Pro Ala Leu Ala Gly His Met Lys Asp Ile Gly Arg Leu Lys Leu		
227	385	390	395
228	Leu Leu Pro Asp Leu Gln Pro Gln Ser Asn Phe Cys Leu Leu Phe Asp		400
229	405	410	415
230	Tyr Arg Leu Ala Gly Asp Lys Val Gly Lys Leu Arg Val Phe Val Lys		
231	420	425	430
232	Asn Ser Asn Asn Ala Leu Ala Trp Glu Lys Thr Thr Ser Glu Asp Glu		
233	435	440	445
234	Lys Trp Lys Thr Gly Lys Ile Gln Leu Tyr Gln Gly Thr Asp Ala Thr		
235	450	455	460
236	Lys Ser Ile Ile Phe Glu Ala Glu Arg Gly Lys Gly Lys Thr Gly Glu		
237	465	470	475
238	Ile Ala Val Asp Gly Val Leu Leu Val Ser Gly Leu Cys Pro Asp Ser		480
239	485	490	495
240	Leu Leu Ser Val Asp Asp Xaa Met Val Leu Ser Leu Tyr Leu Thr Leu		
241	500	505	510
242	Tyr Val Ser Ser Leu Val Phe Leu Ile Leu His His Arg Thr Ser Gly		
243	515	520	525
244	Ile Leu Lys Leu Leu Ala Glu Lys Leu		

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

VERIFICATION SUMMARY  
PATENT APPLICATION US/08/968,800DATE: 04/15/1999  
TIME: 11:14:59

Input Set: H968800.RAW

## Line ? Error/Warning

## Original Text

35 W "N" or "Xaa" used: Feature required  
67 W "N" or "Xaa" used: Feature required  
145 W "N" or "Xaa" used: Feature required  
240 W "N" or "Xaa" used: Feature required  
255 W "N" or "Xaa" used: Feature required  
257 W "N" or "Xaa" used: Feature required  
259 W "N" or "Xaa" used: Feature required  
270 W "N" or "Xaa" used: Feature required  
272 W "N" or "Xaa" used: Feature required  
274 W "N" or "Xaa" used: Feature required  
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315 W "N" or "Xaa" used: Feature required  
330 W "N" or "Xaa" used: Feature required  
347 W "N" or "Xaa" used: Feature required  
361 W "N" or "Xaa" used: Feature required

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Ile Asp Glu Cys Xaa Ser Asn Pro Cys Gln A  
Xaa Xaa Asp Xaa Val Gly Ser Tyr Xaa Cys X  
Thr Gly Lys Xaa Xaa Xaa Cys Glu Xaa Asn  
Xaa Asn Glu Cys Thr Met Xaa Xaa Xaa Cys G  
Val Asn Thr Xaa Gly Ser Tyr Xaa Cys Lys C  
Gly Xaa Xaa Leu Xaa Cys Asp  
Val Xaa Glu Cys Xaa Ser Gly Xaa Gln Xaa X  
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Trp Xaa Pro Xaa Pro Gly Xaa Pro Asn Xaa X  
Cys Val Val Gly Tyr Ile Xaa Xaa Gly G  
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